

SEQUENCE LISTING

<110> Huse, William D.

Freedman, Michael H.

<120> Method for Identifying Optimal Binding Ligands to a
Receptor

<130> P-IX 3280

<140> US 09/169,048

<141> 1998-10-08

<150> 60/112,011

<151> 1997-10-09

<160> 28

<170> PatentIn Ver. 2.0

<210> 1

<211> 24

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (1)..(24)

<400> 1

agc tca agt gta agt ttc atg aac
Ser Ser Ser Val Ser Phe Met Asn

1

5

24

<210> 2

<211> 8

<212> PRT

<213> Mus musculus

<400> 2

Ser Ser Ser Val Ser Phe Met Asn

1

5

<210> 3

<211> 24

<212> DNA

200240-09462360

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 3

agc tca agt gta agg ttc atg aac
Ser Ser Ser Val Arg Phe Met Asn
1 5

24

<210> 4

<211> 8

<212> PRT

<213> Artificial Sequence

<400> 4

Ser Ser Ser Val Arg Phe Met Asn
1 5

<210> 5

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 5

agc gag agt gta aat ctt atg aac
Ser Glu Ser Val Asn Leu Met Asn
1 5

24

<210> 6

<211> 8

<212> PRT

2000-09-14-000000

<213> Artificial Sequence

<400> 6

Ser Glu Ser Val Asn Leu Met Asn
1 5

<210> 7

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 7

agc tca agt gtt aat ttc atg aac
Ser Ser Ser Val Asn Phe Met Asn
1 5

24

<210> 8

<211> 8

<212> PRT

<213> Artificial Sequence

<400> 8

Ser Ser Ser Val Asn Phe Met Asn
1 5

<210> 9

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

ପାତ୍ରବିଦ୍ୟା

<400> 9
agc tca acg gta agt ttc atg aac
Ser Ser Thr Val Ser Phe Met Asn
1 5

24

<210> 10
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 10
Ser Ser Thr Val Ser Phe Met Asn
1 5

<210> 11
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1) .. (24)

<220>
<223> Description of Artificial Sequence: synthetic
construct

<400> 11
agc tca agt gta gcg tat atg aac
Ser Ser Ser Val Ala Tyr Met Asn
1 5

24

<210> 12
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 12
Ser Ser Ser Val Ala Tyr Met Asn
1 5

<210> 13
<211> 24

```
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(24)

<220>
<223> Description of Artificial Sequence: synthetic
      construct
```

<400> 13
agc cag agt gct aag cat atg aac
Ser Gln Ser Ala Lys His Met Asn
1 5

24

<210> 14
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 14
Ser Gln Ser Ala Lys His Met Asn
1 5

```
<210> 15
<211> 24
<212> DNA
<213> Mus musculus
```

<220>
<221> CDS
<222> (1)..(24)

<400> 15
gcc aca tcc aat ttg gct tct gga
Ala Thr Ser Asn Leu Ala Ser Gly
1 5

24

<210> 16
<211> 8
<212> PRT
<213> *Mus musculus*

<400> 16

Ala Thr Ser Asn Leu Ala Ser Gly

1

5

<210> 17

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 17

gcc aca gag aag ttg gct tct gga

Ala Thr Glu Lys Leu Ala Ser Gly

1

5

24

<210> 18

<211> 8

<212> PRT

<213> Artificial Sequence

<400> 18

Ala Thr Glu Lys Leu Ala Ser Gly

1

5

<210> 19

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)..(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 19

gcc aca gtt aat ttg gct tct gga

24

Ala Thr Val Asn Leu Ala Ser Gly
1 5

<210> 20
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 20
Ala Thr Val Asn Leu Ala Ser Gly
1 5

<210> 21
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(24)

<220>
<223> Description of Artificial Sequence: synthetic
construct

<400> 21
gcc aca gtg aat ttg gct tct gga
Ala Thr Val Asn Leu Ala Ser Gly
1 5

24

<210> 22
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 22
Ala Thr Val Asn Leu Ala Ser Gly
1 5

<210> 23
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(24)

<220>
<223> Description of Artificial Sequence: synthetic
construct

<400> 23
gcc aca tcc agg gcg gct tct gga
Ala Thr Ser Arg Ala Ala Ser Gly
1 5

24

<210> 24
<211> 8
<212> PRT
<213> Artificial Sequence

<400> 24
Ala Thr Ser Arg Ala Ala Ser Gly
1 5

<210> 25
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(24)

<220>
<223> Description of Artificial Sequence: synthetic
construct

<400> 25
gcc aca cag aat ttg gct tct gga
Ala Thr Gln Asn Leu Ala Ser Gly
1 5

24

<210> 26
<211> 8
<212> PRT
<213> Artificial Sequence

09639460-DNA0001
<400> 26

Ala Thr Gln Asn Leu Ala Ser Gly
1 5

<210> 27

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<221> CDS

<222> (1)...(24)

<220>

<223> Description of Artificial Sequence: synthetic
construct

<400> 27

gcc aca tcc aat ttg gct tct gga
Ala Thr Ser Asn Leu Ala Ser Gly
1 5

24

<210> 28

<211> 8

<212> PRT

<213> Artificial Sequence

<400> 28

Ala Thr Ser Asn Leu Ala Ser Gly
1 5